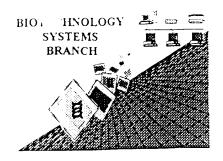
RAW SEQUENCE LISTING ERROR REPORT



The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number:	09/849,526
Source:	OIPE
Date Processed by STIC:	7/24/2001

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,

2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216. PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax) PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE SEE BELOW

Checker Version 3.0

The Checker Version 3.0 application is a state-of the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 – 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST 25

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO). Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money

Checker Version 3.0 can be down loaded from the USPTO website at the following address: http://www.uspto.gov/web/offices/pac/checker

Raw Sequence Listing Error Summary

ERROR DETECTED	SUGGESTED CORRECTION SERIAL NUMBER: 17/54/ 520
ATTN: NEW RULES CASE	S: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE
1Wrapped Nucleics Wrapped Aminos	The number text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to 3; this will prevent "wrapping."
2Invalid Line Length	The rules require that a line not exceed 72 characters in length. This includes white spaces.
3Misaligned Amino Numbering	The numbering under each 5 th amino acid is misaligned. Do not use tab codes between numbers; use space characters, instead.
4Non-ASCII	The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.
5Variable Length	Sequence(s) contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.
6PatentIn 2.0 "bug"	A "bug" in Patentln version 2.0 has caused the <220>-<223 section to be missing from amino acid sequences(s) Normally, Patentln would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.
7Skipped Sequences (OLD RULES)	Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence: (2) INFORMATION FOR SEQ ID NO.X: (insert SEQ ID NO where "X" is shown) (i) SEQUENCE CHARACTERISTICS (Do not insert any subheadings under this heading) (xi) SEQUENCE DESCRIPTION:SEQ ID NO X: (insert SEQ ID NO where "X" is shown) This sequence is intentionally skipped
	Please also adjust the "(ii) NUMBER OF SEQUENCES." response to include the skipped sequences.
8 Skipped Sequences (NEW RULES)	Sequence(s) missing. If Intentional, please insert the following lines for each skipped sequence. <210> sequence id number <400> sequence id number 000
9Use of n's or Naa's (NEW RULES)	Use of n's and/or Xaa's have been detected in the Sequence Listing. Per 1-823 of Sequence Rules, use of < 220 >= 223 \(\) is MANDATORY if n's or Xaa's are present. In < 220 > to < 223 > section, please explain location of n or Xaa, and which residue n or Xaa represents.
10Invalid <213 \cdot Kesponse	Per 1.823 of Sequence Rules, the only valid < 2135 responses are: Unknown, Artificial Sequence, or scientific name (Genus species) > 2205 × 223 section is required when < 2135 response is Unknown or is Artificial Sequence
11Use of ~220 ·	Sequence(s) missing the <220 : "Feature" and a sociated numeric identifiers and responses. Use of <220 > to <223 > is MANDATORY if < 213 \times "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in < 220 \times to < 223 \times section (See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (See 1.823 of Sequence Rules)
12 Patentin 2 0 "bug"	Please do not use "Copy to Disk" function of Patentln version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.
13Misuse of n	n can only be used to represent a single nucleotide in a nucleic acid sequence. N is not used to represent any value not specifically a nucleotide.

AMC/MH - Biotechnology Systems Branch 08/21/2001

" my you are any of the folds 221, 222 or 223, these toolds must be provided by the 220 field. The 221 field on made field for the introduction of the different "features" described in 221, 222 or 223

The types of errors shown exist throughout the Sequence Listing. Please check subsequent sequences for similar errors.

DAIE: 07/24/2001

TIME: 09:10:28

OIPE

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                     Output Set: N:\CRF3\07242001\1849526.raw
      1 < 110 > APPLICANT: Ruff, Thomas G.
              Shukla, Hridayabhiranjan
      1 < 120 - TITLE OF INVENTION: Nucleic Acid Molecules And Other Molecules Associated With
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      ' <130 - FILE REFERENCE: 38-21(51930)B
C--> 9 <140 > CURRENT APPLICATION NUMBER: US/09/849,526
C--> 9 <141> CURRENT FILING DATE: 2001-05-07
      9 <150 PRIOR APPLICATION NUMBER: US 60/202,214
     11 <151 - PRIOR FILING DATE: 2000-05-08
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     14 <151 - PRIOR FILING DATE: 2000-09-05
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     19 < 150 + PRIOR APPLICATION NUMBER: US 09/816,660
     20 <151 - PRIOF FILING DATE: 2001-03-26
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     29 + 223 + \texttt{OTHER} INFORMATION clone ID uc-zmroteosinte002d05b1
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     35 oqqq"toqtq qqtqaqoqqa toqaqatqqo qqoqtoqqat qttqaqtaco qotqottoqt.
                                                                             120
                                                                            180
     37 eggeggeete geetgggeea eggaegaeea etceeteeae aacgeettea geacctaegg
     39 eqaqqteete qaqtenaaqa teateetega teqqqaqaeg caqaqqteec geqqettegq
                                                                             240
     41 etteqteace ttetecaeqq aqqaqqeqat qeqqaaeqee ateqaqqqea tqaaeqqeaa
                                                                             300
     43 quaditiqqac qqicoqcaaba toacoqtoaa oqaqqibobaq toocqoqqoq qooqtiqqaqq 360
     4) namh manan ag filinggig giggeegigg aggeggeg taeggeggig gegggeeg 420
     4° tgatugegge ggeggetaeg geggtggegg egge
                                                                             454
                                Read field 2xC com medica
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     67 ticaqaaqta tattqqtqaq qqctcccqqa tqqttaqqqa actctttqtt atqqccaqqq
     65 aacatgcace atccattata titatqqatq aaattqactc tatcqqatct qctagaatqq
                                                                             240
       uqtotggaac tqqcaacqgt gacagtgaaq ttcaqcgtac tatqcttqaa cttctaaacc
                                                                             300
                                                                             360
     69 agetegatog ttitgaagea teaaacaaaa ttaaggttit gatogeaacg aacagaatag
     7: adatittqqa tdaaqoodtt otgagqootq qooqdataga daqqaagatt qaatttodaa
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RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/849,526

The types of errors shown exist throughout the Sequence Listing. Please check subsequent sequences for similar errors.

RAW SEQUENCE LISTING

DATE: 07/24/2001

PATENT APPLICATION: US/09/849,526 TIME: 09:10 28

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Output Set: N:\CRF3\07242001\1849526.raw

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89 qayactigaac caaaqciiit chaiyacqay qaqraqaaqq qacttaaaca acaccaaqag	18:)								
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113 treattetera gagattigeaa erageraane aggiggaara gaateggage tacaagaceg	180								
115 queaqttgqt geqtaaatea tacaaaaaaa aactaaaaaa ceetgttett attitgctac	$\frac{100}{240}$								
117 tacatecagt eggetacgea aaaaacteec atcaaacete ttattaaaca attteacaca	3(11)								
119 chartatggag acctgoagge tgetcoagtg geoagtetee acteceaaaa actgtaagaa	31:11								
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143 at latigaag agggtagtac aattactica tititototat tacaagaigg tgatticott	240 3(0)								
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163 ccaqcaggte geogaecagg actacqceta aqtaqtgete egiteqtetg gattqagete	180								
165 at ccaacgea gccgcqcqtt tcqatcqgcq tcqtcatttg ctt.tgctctt catttcatca	240								
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	teaggatque aanggeeana teattatett accetecace acctecettt taccgaetqt	3 () ()								
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	- danagaggan bagaggaas bagscoacsy gagseggasg gttgggscoc thotgagta	300								
269		360								
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		430								
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	cottotoayg cottoutogo gtoottogoo ggoyacyato accgacaagg atggaaagaa	180								
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	aqaaqaagot ottiglagic tacataloag qigaagaiga gqottocagi agittigqaac	3 () ()								
297		36O								
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RAW SEQUENCE LISTING

DATE 07/24/2001

PATENT APPLICATION: US/09/849,526 TIME 09:10:28

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		<2.12 · LOCATION: (1)(548)									
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	399 402 403 404 405 407 409 411 1.2 415 427 421 423 425	Attactiq (210 - SEQ ID NO: 15 (211 - LENGTH: 50) (212 - TYPE DNA) (213 - ORGANISM: Lea mays subsp. mexicana) (223 - OTHEE INFORMATION: Clone ID: uC-zmroteosinte008g09b1 (400) SEQUENCE: 15 (cotototoca totocotgot cotococofo oftotidato najnoiajoo ayyyatogoo atgitogaa, t oggaqqagoa coacttogaa togaaggoog acgooggago gtocaagaco taccoqocaqo aggotgoca ogtocgtaag aacggottoa togtoaaqqti togaaqqtto tacctocaaag actggtaago atggccatgo caaatgccac titqttqoca tagacatatt caacggaaa aagottgaag atattgttoo ttoatcacac aactgtqata tocqocatgt gaaccgtact gagtaccago tgattqatat atcaqaqgat qatttqfaa qoottottac ttoaqatqqc aacactaaqq atgatettaq actoccaact qatgagaeto ttqtqqcca qatcaaggaa qqgtttgaaa goqqcaaqqa tottqttgtg	60 120 180 240 360 420 480								
	399 402 403 404 405 407 409 411 1.2 415 427 421 423 425 427	Attactiq (210 - SEQ ID NO: 15 (211 - LENGTH: 50) (212 - TYPE: DNA) (213 - ORGANISM: Lea mays subsp. mexicana) (223 - OTHEE INFORMATION: Clone ID: uC-zmroteosinte008g09b1 (400) - SEQUENCE: 15 (cototototoa totocotqot cotococoto offotoaaq acqueqqaq acqueqqaq aqqotqqaa coacttoqaa toqaaqqooq acqueqqaq aqqotqqaa coqottoqaa toqaaqqooq acqueqqaq aqqotqqaa toqaaqqtta toqaaqqtta tacotoaaaq actggtaaq atggccatqo caaatqocac titqttqooqa taqacatatt caacqqqaaa aaqottqaaq atqqttqata toqqaaqqt qaacqqtot qaqtacaqo tqattqtaa accacaaq actgttaa qoottottaa tteaqatqo aacactaaqq atqatottaq actocoaact qatqaqatot ttqtqqocca qatcaaqqaa qqqttqaaa qooqcaaqqa tottqttqtq actgtcoaat ctqctatqqq qqaqqaq	548 60 120 180 240 360 420								
	399 402 403 404 405 407 409 411 1.2 415 427 425 427 430	Attacttq (210 - SEQ ID NO: 15 (211 - LENGTH: 50? (212 - TYPE - DNA) (213 - ORGANISM: Lea mays subsp. mexicana) (223 - OTHEE INFORMATION: Clone ID: uC-zmroteosinte008g09b1 (400) - SEQUENCE: 15 detectation tetrocotget detectate effectionate majoriages asygnatographic adjoinance adjoinanc	60 120 180 240 360 420 480								
	349 402 403 404 405 407 409 411 1.2 415 427 421 423 425 427 430 431	attacttq c210 - SEQ ID NO: 15 c211 - LENGTH: 50? c212 - TYPE: DNA c213 - ORGANISM: Dea mays subsp. mexicana c223 - OTHEE INFORMATION: Cione ID: uC-zmroteosinte008g09b1 c400: SEQUENCE: 15 ceteteteca tetecetget cetecacete ettettaate najnemajes aggyategee stattogrant eggaggage ecaettegaa tegaaggeeg aegeeggage gtocaagaee taccequage aggetgeae egteegtaag aaeggettea tegteateaa gaacegeece tacaaggtig tggaggtte taceteaaag aetggtaage atggeeatge caaatgeeae titgtitgeea tagacatatt caacggaaa aagettgaag atattgttee ticateaaae aactgtgata ticcquatgt gaacegtact gagtaceage tgattgatat atcaqaagat gattigtaa geettettac ticaqatgge aacactaagg atgatettag aeteccaact gatgagacte tigtggeeae qateaaggaa gggttigaaa geggeaagga tettgttgt aetgiceadt etgetatggg ggaggag 210: SEQ ID NO 16 c211: LENGTH: 559	60 120 180 240 360 420 480								
	349 402 403 404 405 407 409 411 1.2 415 427 421 423 427 430 431 432	attacttq 219 - SEQ ID NO: 15 211 - LENGTH: 50° 212 - TYPE: DNA 213 - ORGANISM: Dea mays subsp. mexicana 223 - OTHEE INFORMATION: Cione ID: uC-zmroteosinte008g09b1 400: SEQUENCE: 15 ceteteteca tetecetget cetecoccic citetidate najoriages agguategec tacceopage aggetqeac egiceqtaag aacggetea teqteateaa gaacegeece tacacequage aggetqeac egiceqtaag aacggetea teqteateaa gaacegeece tacacequage tegaaggaaa aacggetaa atggeaatge caaatgceac titqtiqeea tagacatat caaceggaaa aacggtaage atggeaatge caaatgceac titqtiqeea tagacatat caaceggaaa aacgtaaga atatigtice ticatcacac aactutqata ticcqcatgt gaacegtact gagtaccage tgatigatat atcaqaaggat gattigata titqtqqa qeetictata ticaqatqqa aacactaaqa atgatettaa acteccaact gatgagacte tiqtqqeeca qatcaaggaa qggtitgaaa geggcaaqqa tettqttgtg actgiccaat etqetatgqq qqagqag 210: SEQ ID NO 16 211: LENGTH: 559 212: TYPE: DNA	60 120 180 240 360 420 480								
	349 402 403 404 405 407 409 411 1.2 415 427 421 423 421 423 431 432 433	attacttq (210 - SEQ ID NO: 15 (211) LENGTH: 50° (212) TYPE: DNA (213) ORGANISM: Dea mays subsp. mexicana (223) OTHEE INFORMATION: Cione ID: uc-zmroteosinte008g09b1 (400) SEQUENCE: 15 cototototoa totocotgot cotocococc ottotidate najneragos aggadose taccoquago aggotagoa coacttogaa togaaggoog aegocagago gtocaagaco taccoquago aggotagoa ogtocquaa aeggottoa togtocatoa gaacogococ tacaaqqtig tagaaqqtto tacctoaaaa actgataago atggocatgo caaatgocac titqtiqoca tagacatatt caacaggaaa aaggottaa togtocatgo caaatgocac titqtiqoca tagacatatt caacaggaaa aagettgaag atatigttoo ticatcacac aactgiqata ticoqocatgi gaaccgtact gagtaccago tgattqatat atcaqaqqat qaatitqiqaa qocttottac ticaqatqqc aacactaaqq atgatottaq actoccaact qatgagacte tiqqqqocoa qatcaaggaa qggtitgaaa goqqcaaqqa tottqttqq qqaqqaq (210) SEQ ID NO 16 (211) LENGTH: 559 (212) TYPE: DNA (213) ORGANISM Dea mays subsp. mexicana	60 120 180 240 360 420 480								
	349 402 403 404 405 407 409 411 1.2 415 427 421 423 421 423 431 432 433	attacttq 219 - SEQ ID NO: 15 211 - LENGTH: 50° 212 - TYPE: DNA 213 - ORGANISM: Dea mays subsp. mexicana 223 - OTHEE INFORMATION: Cione ID: uC-zmroteosinte008g09b1 400: SEQUENCE: 15 ceteteteca tetecetget cetecoccic citetidate najoriages agguategec tacceopage aggetqeac egiceqtaag aacggetea teqteateaa gaacegeece tacacequage aggetqeac egiceqtaag aacggetea teqteateaa gaacegeece tacacequage tegaaggaaa aacggetaa atggeaatge caaatgceac titqtiqeea tagacatat caaceggaaa aacggtaage atggeaatge caaatgceac titqtiqeea tagacatat caaceggaaa aacgtaaga atatigtice ticatcacac aactutqata ticcqcatgt gaacegtact gagtaccage tgatigatat atcaqaaggat gattigata titqtqqa qeetictata ticaqatqqa aacactaaqa atgatettaa acteccaact gatgagacte tiqtqqeeca qatcaaggaa qggtitgaaa geggcaaqqa tettqttgtg actgiccaat etqetatgqq qqagqag 210: SEQ ID NO 16 211: LENGTH: 559 212: TYPE: DNA	60 120 180 240 360 420 480								
	349 402 403 404 405 407 409 411 1.2 415 427 421 423 421 433 431 432 433 435	attacttq (210 - SEQ ID NO: 15 (211) LENGTH: 50° (212) TYPE: DNA (213) ORGANISM: Dea mays subsp. mexicana (223) OTHEE INFORMATION: Cione ID: uc-zmroteosinte008g09b1 (400) SEQUENCE: 15 cototototoa totocotgot cotocococc ottotidate najneragos aggadose taccoquago aggotagoa coacttogaa togaaggoog aegocagago gtocaagaco taccoquago aggotagoa ogtocquaa aeggottoa togtocatoa gaacogococ tacaaqqtig tagaaqqtto tacctoaaaa actgataago atggocatgo caaatgocac titqtiqoca tagacatatt caacaggaaa aaggottaa togtocatgo caaatgocac titqtiqoca tagacatatt caacaggaaa aagettgaag atatigttoo ticatcacac aactgiqata ticoqocatgi gaaccgtact gagtaccago tgattqatat atcaqaqqat qaatitqiqaa qocttottac ticaqatqqc aacactaaqq atgatottaq actoccaact qatgagacte tiqqqqocoa qatcaaggaa qggtitgaaa goqqcaaqqa tottqttqq qqaqqaq (210) SEQ ID NO 16 (211) LENGTH: 559 (212) TYPE: DNA (213) ORGANISM Dea mays subsp. mexicana	60 120 180 240 360 420 480								
	349 402 403 404 405 407 409 411 1.2 415 427 421 423 421 433 431 432 433 435 436	Attacttq (210 - SEQ ID NO: 15 (211 - LENGTH: 5007 (212 - TYPE: DNA (213 - ORGANISM: Mea mays subsp. mexicana (223 - OTHEE INFORMATION: Cione ID: uC-zmroteosinte008g09b1 (400) - SEQUENCE: 15 ceteteteca tetecetqet cetecocce ettettaate majoriagos aggadiogee (digtegga t eggaggagea ecacttegaa tegaaggeeg aegeeggage gtecaagaee taceegeage aggetigeae egteegtaag aeggetiea tegteateaa gaacegeeee tacaaggif tiggaggitte taceteaaag aetggtaage atggeeatge caaatgeeae (ittgitigeea tagacatatt caaceggaaa aagettgaag atatigtee ticateacae (aactgigata ticegeatgt gaacegtact gagtaceage tgattgatat atcagaggat (gattigtaa geettettae ticaqatqqe aacactaagg atgatettag aeteceaact (atgagaete tigtigeeea gateaaggaa gggttigaaa geeggeaagga tettgitegiceaat ciqetatggg gaaggag (210: SEQ ID NO 16 (211 LENGTH: 559 (212 TYPE: DNA (213 ORGANISM Cea mays subsp. mexicana	60 120 180 240 360 420 480								

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/849,526

DATE: 07/24/2001 TIME: 09:10:29

Input: Set. : D:\Pa_00348.rpt

Output Set: N:\CRF3\07242001\1849526.raw

L:9 M:270 C: Current Appl.cation Number differs, Replaced Current Application No L:9 M:271 C: Current Filing Date differs, Replaced Current Filing Date 1.395 M 341 W: (46) "n" or "Xaa" used, for SEQ ID#:14 I, 457 M. 341 W: (46) "n" or "Xaa" used, for SEQ ID#:16 I...1001 M·341 W. (46) "n" or "Xaa" used, for SEQ II)#.48 L 11'5 M:341 W (46) "n" or "Xaa" used, for SEQ ID# 44 L 1349 M.341 W. (46) "n" or "Xaa" used, for SEQ ID#:51 I::1351 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:51 L:1353 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:51 L::1493 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:57 I. 1533 M.341 W. (46) "n" or "Xaa" used, for SEQ ID#:58 I. 2033 M.341 W: (46) "n" or "Xaa" used, for SEQ ID#:81 L 2151 M.341 W: (46) "n" or "Xaa" used, for SEQ ID#:84 I. 2229 M:341 W: (46) "n" or "Xaa" used, for SEQ IF#.87 L.2257 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#.88 L:2287 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:89 L:2349 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:91 L.2531 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:98 L:2965 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:116 I. 3065 M:341 W: (46) "n" or "Xax" used, for SEQ ID#:130 I. 3067 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:120 L:3289 M 341 W: (46) "n" or "Xaa" used, for SEQ ID#:129 L:3409 M:341 W. (46) "n" or "Xaa" used, for SEQ ID#:154 L:3543 M:341 W. (4b) "n" or "Xaa" used, for SEQ II#:159 L:3679 M 341 W: (46) "n" or "Xia" used, for SEQ II:#.144 (4b) "n" or "Xaa" used, for SEQ ID# 150 L:3823 M:341 W I..4119 M.341 W: (46) "n" or "Xaa" used, for SEQ II# 162 L 4201 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#.165 L 4395 M.341 W: (46) "n" or "Xaa" used, for SEQ ID#:1"3 I. 4405 M 341 W: (46) "n" or "Xaa" used, for SEQ ID#:177 L.4645 M.341 W. (46) "n" or "Xaa" used, for SEQ ID#:183 L.4969 M.341 W. (46) "n" or "Xaa" used, for SEQ ID#:187 L 5001 M.341 W. (46) "n" or "Xaa" used, for SEQ ID#:148 (46) "n" or "Xaa" used, for SEQ ID# 202 L.5103 M.341 W L blos Mr341 Wr (46) "n" or "Xaa" used, for SEQ ID#.204 L.5587 M 341 W: (4b) "n" or "Xaa" used, for SEQ ID# 220 [1.5589 M.341 W: (46) "n" or "Xaa" used, for SEQ ID#:220 L 5939 M·341 W· (46) "n" or "Xaa" used, for SEQ ID#.236 I. 6125 M 341 W: (46) "n" or "Xaa" used, for SEQ ID# 244 (4b) "n" or "Xaa" used, for SEQ ID# 200 L 6495 M 341 W L 6497 M:341 W. (46) "n" or "Xaa" used, for SEQ ID#.260 L.6601 M 341 W: (46) "n" or "Xaa" used, for SEQ ID#.264 (46) "n" or "Xaa" used, for SEQ ID# 276L 6859 M 341 W L 7259 M:341 W (46) "n" or "Xaa" used, for SEQ ID# 291 (46) "n" or "Xaa" used, for SEQ ID# 294 L 7335 M 341 W (46) "n" or "Xaa" used, for SEQ ID#:296 L:7381 M 341 W L:7411 M 341 W. (46) "n" or "Xaa" used, for SEQ ID#:297 L:7425 M:341 W (46) "n" or "Xaa" used, for SEQ ID# 297

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/849,526 TIME: 09:10 29

DATE: 07/24/2001

Input Set : D:\Pa_00348.rpt

Output Set: N:\CRF3\07242001\1849526.raw

L: 7441	М	341	W:	(46)	" n "	$\circ r$	"Xaa"	used,	for	SEQ	11)#	298
L: 7455	1-1	341	W_{\perp}	(46)	" n "	or	"Xaa"	used,	for	SEQ	[]]#	298
1.:7551	1:1	341	W:	(46)	$^{\rm n}{\rm n}^{\rm n}$	$\circ r$	"Xaa"	used,	for	SEQ]])#	302
L:7619	Μ	: 341	W:	(46)	" II "	or	"Xaa"	used,	for	SEQ	I [] #	305